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<110> SUNTORY LIMITED

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<151> 1998-04-01

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<170> PatentIn Ver. 2.0

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Asn Lys

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09-10-2017

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gtc aat tct tta ctt act aag gta gaa act gcc tta aaa gaa gct tct 150

30 35 40

Gly Ser Asn Glu Ala Ala Leu Glu Ala Leu Glu Gly Leu Lys Gly Glu

45 50 55

atc cag aca aaa cca gac cga gtt gga caa gcc aca aaa atc ctt gga 246

Ile Gln Thr Lys Pro Asp Arg Val Gly Gln Ala Thr Lys Ile Leu Gly

20 60 65 70

tct gtc gga tca gct cta gga aaa tta aat tct gga gat gca acc aaa 294

Ser Val Gly Ser Ala Leu Gly Lys Leu Asn Ser Gly Asp Ala Thr Lys

75 80 85

atc att tct ggt tgc ctc gac att gtt gca gga att gca aca act ttt 342

25 Ile Ile Ser Gly Cys Leu Asp Ile Val Ala Gly Ile Ala Thr Thr Phe

90 95 100 105

gga ggc cct gtc ggg atg gga atc gga gcc gta gct tct ttt gtt tct 390

Gly Gly Pro Val Gly Met Gly Ile Gly Ala Val Ala Ser Phe Val Ser

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tca att cta tca ttg ttt act gga agc tca gca aag aac tca gtt gct 438
 Ser Ile Leu Ser Leu Phe Thr Gly Ser Ser Ala Lys Asn Ser Val Ala
 125 130 135
 gcc gtt att gat aga gct tta agc aag cat cgc gat gag gcc atc caa 486
 5 Ala Val Ile Asp Arg Ala Leu Ser Lys His Arg Asp Glu Ala Ile Gln
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 aga cat gca gca ggt gcc aag aga gat ttt gct gaa tca tct gca ttc 534
 Arg His Ala Ala Gly Ala Lys Arg Asp Phe Ala Glu Ser Ser Ala Phe
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 Ile Gln Val Met Lys Gln Gln Ser Asn Leu Thr Asp Ser Asp Leu Ser
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 15 190 195 200
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 25 gga aat gca gaa cac gtg gca agt gct gtg gaa aac gct aat agg gta 822
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 250 255 260 265
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 Asn Lys Glu Leu Ala Ala Asp Thr Leu Asp Phe Leu His Lys Leu Ile

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	Pro Glu Gln Ala Leu Ile Gly Ala Val Tyr His Pro Ile Ser Ala Ser			
	285	290	295	
5	gaa act agc aaa gca ata tta aat tac acg aaa tac ttt gga gtt cca			966
	Glu Thr Ser Lys Ala Ile Leu Asn Tyr Thr Lys Tyr Phe Gly Val Pro			
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	gat gtt ccc cgt cct att gga aac cgc aga tac aaa ttt aca aat agt			1014
	Asp Val Pro Arg Pro Ile Gly Asn Arg Arg Tyr Lys Phe Thr Asn Ser			
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	Tyr Trp Asn Thr Tyr Ser Ile Cys Ser Glu Ala Tyr Met Gly Asn Tyr			
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	atg ttc aga ggc tgt tct aac gtt cgg aat cca aat atc agg gta tcc			1110
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	aaa atg tct gat ggg ttt tac acc atg gag aat agc gat cgg agg aag			1158
	Lys Met Ser Asp Gly Phe Tyr Thr Met Glu Asn Ser Asp Arg Arg Lys			
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20	ttg tat atc acc aag cat gac caa gga tgg gga tgg ggt act ttg gat			1206
	Leu Tyr Ile Thr Lys His Asp Gln Gly Trp Gly Trp Gly Thr Leu Asp			
	380	385	390	
	gag gat cca ggt gac caa ggc cat atg agg ttc att cct ttg aga cat			1254
	Glu Asp Pro Gly Asp Gln Gly His Met Arg Phe Ile Pro Leu Arg His			
25	395	400	405	
	ggg aag tat atg gta agc tct aag agg tgg ccc aac tgg ttc atg tat			1302
	Gly Lys Tyr Met Val Ser Ser Lys Arg Trp Pro Asn Trp Phe Met Tyr			
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Met Glu Ser Ser Ala Ser Gly Tyr Ile Arg Ser Trp Glu Asn Asn Pro

430

435

440

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Gly Pro Gln Gly His Trp Ser Ile Thr

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445

450

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<213> CARYBDEA RASTONII

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20

25

30

Val Glu Thr Ala Leu Lys Glu Ala Ser Gly Ser Asn Glu Ala Ala Leu

35

40

45

Glu Ala Leu Glu Gly Leu Lys Gly Glu Ile Gln Thr Lys Pro Asp Arg

50

55

60

25

Val Gly Gln Ala Thr Lys Ile Leu Gly Ser Val Gly Ser Ala Leu Gly

65

70

75

80

Lys Leu Asn Ser Gly Asp Ala Thr Lys Ile Ile Ser Gly Cys Leu Asp

85

90

95

Ile Val Ala Gly Ile Ala Thr Thr Phe Gly Gly Pro Val Gly Met Gly

	100	105	110	
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	115	120	125	
	Gly Ser Ser Ala Lys Asn Ser Val Ala Ala Val Ile Asp Arg Ala Leu			
5	130	135	140	
	Ser Lys His Arg Asp Glu Ala Ile Gln Arg His Ala Ala Gly Ala Lys			
	145	150	155	160
	Arg Asp Phe Ala Glu Ser Ser Ala Phe Ile Gln Val Met Lys Gln Gln			
	165	170	175	
10	Ser Asn Leu Thr Asp Ser Asp Leu Ser Ile Ile Ala Ala Asn Val Pro			
	180	185	190	
	Val Tyr Lys Phe Ser Asn Phe Ile Gly Gln Leu Glu Ser Arg Ile Ser			
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	Gln Gly Ala Ala Thr Thr Ser Leu Ser Asp Ala Lys Arg Ala Val Asp			
15	210	215	220	
	Phe Ile Leu Leu Tyr Cys Gln Leu Val Val Met Arg Glu Thr Leu Leu			
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	Val Asp Leu Ala Ile Leu Tyr Arg Lys Gly Asn Ala Glu His Val Ala			
	245	250	255	
20	Ser Ala Val Glu Asn Ala Asn Arg Val Asn Lys Glu Leu Ala Ala Asp			
	260	265	270	
	Thr Leu Asp Phe Leu His Lys Leu Ile Pro Glu Gln Ala Leu Ile Gly			
	275	280	285	
	Ala Val Tyr His Pro Ile Ser Ala Ser Glu Thr Ser Lys Ala Ile Leu			
25	290	295	300	
	Asn Tyr Thr Lys Tyr Phe Gly Val Pro Asp Val Pro Arg Pro Ile Gly			
	305	310	315	320
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Cys Ser Glu Ala Tyr Met Gly Asn Tyr Met Phe Arg Gly Cys Ser Asn

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Val Arg Asn Pro Asn Ile Arg Val Ser Lys Met Ser Asp Gly Phe Tyr

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5 Thr Met Glu Asn Ser Asp Arg Arg Lys Leu Tyr Ile Thr Lys His Asp

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Gln Gly Trp Gly Trp Gly Thr Leu Asp Glu Asp Pro Gly Asp Gln Gly

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390

395

400

His Met Arg Phe Ile Pro Leu Arg His Gly Lys Tyr Met Val Ser Ser

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410

415

Lys Arg Trp Pro Asn Trp Phe Met Tyr Met Glu Ser Ser Ala Ser Gly

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425

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Tyr Ile Arg Ser Trp Glu Asn Asn Pro Gly Pro Gln Gly His Trp Ser

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Ile Thr

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